

E. Willis

Re-run

STC Copy #11

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/522,342DATE: 10/15/2001  
TIME: 15:49:39Input Set : A:\P1219P1-US Sequence Listing.txt  
Output Set: N:\CRF3\10152001\I522342.raw

3 <110> APPLICANT: Stewart, Timothy A.  
 4 Tomlinson, Elizabeth  
 5 Goddard, Audrey  
 6 Gurney, Austin L.  
 8 <120> TITLE OF INVENTION: FIBROBLAST GROWTH FACTOR-19 (FGF-19) NUCLEIC ACIDS AND  
 9 POLYPEPTIDES AND METHODS FOR THE TREATMENT OF OBESITY  
 11 <130> FILE REFERENCE: P1219P1-US  
 13 <140> CURRENT APPLICATION NUMBER: US 09/522,342  
 14 <141> CURRENT FILING DATE: 2000-03-09  
 16 <150> PRIOR APPLICATION NUMBER: US 60/066,840  
 17 <151> PRIOR FILING DATE: 1997-11-25  
 19 <150> PRIOR APPLICATION NUMBER: US 09/158,342  
 20 <151> PRIOR FILING DATE: 1998-09-21  
 22 <150> PRIOR APPLICATION NUMBER: US 09/284,663  
 23 <151> PRIOR FILING DATE: 1999-04-15  
 25 <150> PRIOR APPLICATION NUMBER: PCT/US98/25190  
 26 <151> PRIOR FILING DATE: 1998-11-25  
 28 <160> NUMBER OF SEQ ID NOS: 5  
 30 <210> SEQ ID NO: 1  
 31 <211> LENGTH: 2137  
 32 <212> TYPE: DNA  
 33 <213> ORGANISM: Homo Sapien  
 35 <400> SEQUENCE: 1  
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 40 tgggcggggc caccggcgt gggacaagaa gccggccgct gcctgcccgg 150  
 42 gccccggggag ggggctgggg ctggggccgg aggccgggtg tgagtgggtg 200  
 44 tttgcggggg gggggggctt gatgcaatcc cgataagaaa tgctcgggtg 250  
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 52 gtcggccggcc agcctcccgcc accccccatcg ccggagctgc gccgagagcc 450  
 54 ccaggggaggt gccatcgccgaa gccccgtgtgt ggtggccac gatggatcc 500  
 56 tggccggccct ctggctggcc gtggccggggc gccccctcgcc cttctcgac 550  
 58 gccccggggcc acgtgcacta cggctggggc gaccccatcc gcctgcccgg 600  
 60 cctgtacacc tccggccccc acgggccttc cagctgcttc ctgcgcattcc 650  
 62 gtggccgcgg cgtcgtggac tgcgcgggg gccagagcgc gcacagtttg 700  
 64 ctggagatca aggcaatcg tctgcggacc gtggccatca agggcgtgca 750  
 66 cagcgtgcgg tacctctgca tggggccgca cggcaagatg caggggctgc 800  
 68 ttcaatgttgc ggagggatc tggctttcg aggaggat cccggccat 850  
 70 ggctacaatg tggatcgatc cgagaagcac cgcctccgg tctccctgag 900  
 72 cagtgcacaa cagcggcagc tggatcgatc cagaggctt cttccactct 950  
 74 ctcatttcct gccccatcg cccatggtcc cagaggagcc tgaggaccc 1000  
 76 agggccact tggatcgatc cttccactct tcggccctgg agaccgacag 1050  
 78 catggaccca tttggcttg tcaccggact ggaggccgtg aggagtcac 1100  
 80 gctttgagaa gtaactgaga ccatggccgg gccttccac tgctgcccagg 1150  
 82 ggctgtggta cctgcagcgt gggggacgtg cttctacaag aacagtcctg 1200

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84 agtccacgtt ctgttagct ttaggaagaa acatctagaa gttgtacata 1250  
 86 ttcagagttt tccatggca gtgcagttt ctagccaata gacttgtctg 1300  
 88 atcataacat tgtaagcctg tagcttgcgg agctgctgcc tggggcccca 1350  
 90 ttctgtccc tcgagggtgc tggacaagct gctgcactgt ctcagttctg 1400  
 92 ctgtaaatacc tccatcgatg gggaaactcac ttccttgaa aaaattctta 1450  
 94 tgtcaagctg aaattctcta atttttctc atcaactccc caggagcagc 1500  
 96 cagaagacag gcagtagttt taatttcagg aacaggtgat ccaactctgta 1550  
 98 aaacagcagg taaatttcac tcaacccat gtgggaattt atctataatct 1600  
 100 ctacttccag ggaccatttg cccttccaa atccctccag gccagaactg 1650  
 102 actggagcag gcatggccca ccaggcttca ggagtagggg aaggctggag 1700  
 104 ccccaactcca gcccctggac aacttgagaa ttccccctga ggccagttct 1750  
 106 gtcatggatg ctgtcctgag aataacttgc tgcgtccgggt tcacctgttt 1800  
 108 ccatctccca gcccaccaggc cctctgccc ctcacatgc ctccccatgg 1850  
 110 attggggcct cccaggcccc ccacctttagt tcaacctgca cttcttgcc 1900  
 112 aaaaatcagg aaaagaaaaag atttgaagac cccaaagtctt gtcaataact 1950  
 114 tgctgtgtgg aagcagcggg ggaagaccta gaacccttc cccagcaatt 2000  
 116 gttttccaa catgatattt atgagtaatt tattttgata tgtacatctc 2050  
 118 ttattttctt acattattta tgccccaaa ttatattat gtatgttaagt 2100  
 120 gaggtttgtt ttgtatatta aaatggagtt tgtttgt 2137  
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 123 <211> LENGTH: 216  
 124 <212> TYPE: PRT  
 125 <213> ORGANISM: Homo Sapien  
 127 <400> SEQUENCE: 2  
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 129 1 5 10 15  
 131 Leu Trp Leu Ala Val Ala Gly Arg Pro Leu Ala Phe Ser Asp Ala  
 132 20 25 30  
 134 Gly Pro His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg  
 135 35 40 45  
 137 His Leu Tyr Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu  
 138 50 55 60  
 140 Arg Ile Arg Ala Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser  
 141 65 70 75  
 143 Ala His Ser Leu Leu Glu Ile Lys Ala Val Ala Leu Arg Thr Val  
 144 80 85 90  
 146 Ala Ile Lys Gly Val His Ser Val Arg Tyr Leu Cys Met Gly Ala  
 147 95 100 105  
 149 Asp Gly Lys Met Gln Gly Leu Leu Gln Tyr Ser Glu Glu Asp Cys  
 150 110 115 120  
 152 Ala Phe Glu Glu Ile Arg Pro Asp Gly Tyr Asn Val Tyr Arg  
 153 125 130 135  
 155 Ser Glu Lys His Arg Leu Pro Val Ser Leu Ser Ser Ala Lys Gln  
 156 140 145 150  
 158 Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu Pro Leu Ser His Phe  
 159 155 160 165  
 161 Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro Glu Asp Leu Arg  
 162 170 175 180  
 164 Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu Glu Thr Asp

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165 185 190 195  
167 Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala Val Arg  
168 200 205 210  
170 Ser Pro Ser Phe Glu Lys  
171 215  
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174 <211> LENGTH: 26  
175 <212> TYPE: DNA  
176 <213> ORGANISM: Artificial Sequence  
178 <220> FEATURE:  
179 <223> OTHER INFORMATION: Synthetic oligonucleotide probe  
181 <400> SEQUENCE: 3  
182 atccggccag atggctacaa tgtgta 26  
184 <210> SEQ ID NO: 4  
185 <211> LENGTH: 22  
186 <212> TYPE: DNA  
187 <213> ORGANISM: Artificial Sequence  
189 <220> FEATURE:  
190 <223> OTHER INFORMATION: Synthetic oligonucleotide probe  
192 <400> SEQUENCE: 4  
193 ccagtccgggt gacaagccaa 22  
195 <210> SEQ ID NO: 5  
196 <211> LENGTH: 42  
197 <212> TYPE: DNA  
198 <213> ORGANISM: Artificial Sequence  
200 <220> FEATURE:  
201 <223> OTHER INFORMATION: Synthetic oligonucleotide probe  
203 <400> SEQUENCE: 5  
204 gcctccgggt ctccctgagc agtgccaaac agcggcagtg ta 42

**VERIFICATION SUMMARY**

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